

Appl. No. : 10/786,518
 Filed : February 4, 2004

AMENDMENTS TO THE SPECIFICATION

Please replace Tables 2 through 10, which are located between paragraphs [0061] and [0062] of the specification, with the following paragraphs:

Table 2: CDH23 primers

primer name	primer sequence	SEQ ID NO.	bases	product size	primer pair	exons
CDH23-1F	CACTGTGCTATACCCAGGATAGGACAATGTTA	<u>1</u>	32	6886 bp	1F/1R	2 to 3
CDH23-1R	TCAGGTGGAAGATGACCTAACCTGTAAGATC	<u>2</u>	32			
CDH23-2F	GATACCACATGACACACTGTGACAAGT	<u>3</u>	28	1426 bp	2F/2R	4 to 6
CDH23-2R	GACTCTCACCTACACCATGGTGGTCTG	<u>4</u>	28			
CDH23-3F	TATGTATTCTTACACTAACCTGTGAGATATG	<u>5</u>	33	12362 bp	3F/3R	7 to 9
CDH23-3R	TAGCCCTCAGAGCCTGAGATGCCTACTGGCTC	<u>6</u>	32			
CDH23-4F	TGAGTCTTAATGCCAGAGAGGAG	<u>7</u>	25	2604 bp	4F/4R	10 to 11
CDH23-4R	TGAGATGGAGTCTTACTCTTGTTC	<u>8</u>	25			
CDH23-5F	CCAGAACGCTATGCCCATCAGAGG	<u>9</u>	24	3425 bp	5F/5R	12 to 13
CDH23-5R	GCAACCAAGAGTACTGACAGATACA	<u>10</u>	25			
CDH23-6F1	TGTAGGTAGAAGGCGTGCAGGAGCCAGTCGC	<u>11</u>	34	6878bp	6F/6R	14 to 16
CDH23-6R1	GGTCGAGTGTGGCTGCTCAGCCTCCGAGTAT	<u>12</u>	34			
CDH23-6F1b	CCAAAGGAGACGTGCGAGAGGAACAT	<u>13</u>	26	4601 bp	6F1b/6R1b	14 to 16
CDH23-6R1b	TTCTGAGTAGCCAGAGTGTTCAGG	<u>14</u>	25			
CDH23-7F	ACCTCAGTCGAGATGTTGAGGCTCCAGGTGTT	<u>15</u>	33	13282 bp	7F/7R	17 to 21
CDH23-7R	CTATTGCAAGAGCCAGCTCAGAGGGACACAGA	<u>16</u>	32			
CDH23-8F	GAGGGTTGATGAGGAGGAACCCAGTCTCAA	<u>17</u>	32	12314 bp	8F/8R	22 to 27
CDH23-8R	ATTAACTCGCTGGCTCTAGGATTCAGTAAGAG	<u>18</u>	33			
CDH23-9F	GTAGGATGCGTGAAGGGAAGGAAAGGAAC	<u>19</u>	30	8499 bp	9F/9R	28 to 31
CDH23-9R	GTGCACACAGAAGGAGCTAACCAATGTTGG	<u>20</u>	31			
CDH23-10F	GTTATGCCGGACAGAGGAAGTGACATGGAGGT	<u>21</u>	32	7903 bp	10F/10R	32 to 36
CDH23-10R	CAAGGATTGCGCTGCTGTGTTGAAATTCCATT	<u>22</u>	32			
CDH23-11F	GAGTCACATGGAGTGAGTCAGCCCAGGAGAA	<u>23</u>	32	11691 bp	11F/11R	37 to 43
CDH23-11R	ACAATGACCAAGACTGCTCTTCAACCAACAGAC	<u>24</u>	32			
CDH23-12F3a	TTATGACTTGTCTGATCTTCTTCTGATG	<u>25</u>	32	7912bp	12F3a/12R3a	44 to 46
CDH23-12R3a	TTTGTAAAAGTAAATTACACTACCGACTG	<u>26</u>	32			
CDH23-12F4	ACACAGAGGTGCAAGAGAGGTGACATAACTCC	<u>27</u>	32	6815bp	12F4/12R6	47 to 53
CDH23-12R6	TAGCACAGCCATATAGTAACCACTACCGACTG	<u>28</u>	34			
CDH23-13F	CTTGGACACCCATGATGTCTGGGGGGGGGA	<u>29</u>	32	12462 bp	13F/13R	53 to 68
CDH23-13R	GTGACCCCTCCTAACCTGTGCTTAGATGCTAACATT	<u>30</u>	37			

Table 3: GJB2 primers

primer name	primer sequence	SEQ ID	bases	product size	primer pair	exons
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Appl. No. : 10/786,518
 Filed : February 4, 2004

		NO:				
GJB2-1F	AACCTTAGTCCTGGCACATTGTTGAA	<u>31</u>	27	6478 bp	1F/1R2	1 to 2
GJB2-1R2	AACACCACATTGTCCATAGACTGATATG	<u>32</u>	28			
GJB2-1F2	AGTCAATGCTAATAATGGTGGCAATCACG	<u>33</u>	29	7156 bp	1F2/1R2	2
GJB2-1R2	AACACCACATTGTCCATAGACTGATATG	<u>34</u>	28			

Table 4: GJB6 primers

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
GJB6-1F	TATGAGAAGGCTGGATCACCCAGAAAGACTG	<u>35</u>	31	11,112 bp	1F/1R	all 4 exons
GJB6-1R	TGAGGACATCATCCTAGTGTGTCGACAAGTGG	<u>36</u>	31			
GJB6-2F-1	TGTGTTCTGGATTAATGCAAACAGC	<u>37</u>	26	2361 bp	2F-1/2R-2	all 4 exons
GJB6-2R-2	GGACATCATCCTAGTGTGTCGACAAGT	<u>38</u>	26			
GJB6-2F-2	AGCCAATCTGGGTAAATGGATCAGAC	<u>39</u>	26	2383 bp	2F-2/2R-1	all 4 exons
GJB6-2R-1	AGTGCTCTGTAGGCTGCTAAACTTAG	<u>40</u>	26			

Table 5: KCNE1 primers

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
KCNE-1F	GAAAGAGGCATGGAGAGTGAT	<u>41</u>	21	1719 bp	1F/1R1	1 to 2
KCNE-1R1	CTGAAGCTCACTGACGTCTGT	<u>42</u>	20			
KCNE-1F1	CATGGATACCAAGAGACAAC	<u>43</u>	21	1724 bp	1F1/1R	
KCNE-1R	AGGATCACCTCCATTGATTC	<u>44</u>	20			
KCNE-2F	TCCATTAAGGAAGGACCTG	<u>45</u>	20	437bp	2F/2R	3
KCNE-2R	TAAACATTCAAGCGAATGCAG	<u>46</u>	20			
KCNE-3F1	AACCAGTCTGACTAGTCTGCATAAGCT	<u>47</u>	28	4893 bp	3F1/3R2	4
KCNE-3R2	GAGTCTGTTATGCTTGTCAAGGTGT	<u>48</u>	28			

Table 6: KCNQ1 primers

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
KNQ1-1F1	GGTAAATGCACACTGGAACG	<u>49</u>	20	1168bp	1F1/1R1	1
KNQ1-1R1	AGGATTACACACCTGGACTAC	<u>50</u>	20			
KNQ1-2F	ATCCACGTGGCAGCATGTGTTG	<u>51</u>	22	564bp	2F/2R	2
KNQ1-2R	CTTTCAGACCAACCAGCTCCAGGTT	<u>52</u>	24			
KNQ1-3F	ATGAGCTGAAGCTGCTAGCCTTC	<u>53</u>	24	2709bp	3F/3R	3 to 6
KNQ1-3R	TCCAAGCACAGGTTGTGGACAG	<u>54</u>	23			
KNQ1-4F	GCTCTGTTCTGGTCTTCCGGAGT	<u>55</u>	27	5779bp	4F/4R1	7 to 10
KNQ1-4R1	GACAGGTCTGCCATCCAATCGTCAGGT	<u>56</u>	27	6183 bp		
KNQ1-5F1	GACACTGAGGTGTCAGGCACCT	<u>57</u>	22	532bp	5F1/5R1	11
KNQ1-5R1	AGGATCATGTTCCCAGGCTCA	<u>58</u>	21			

Appl. No. : 10/786,518
 Filed : February 4, 2004

KNQ1-6F	TTGCTATGGCTGCCATGTGTCAGCAGCATAG	<u>59</u>	31	9883bp	6F/6R	12 to 15
KNQ1-6R	TCTGCCACCCCTCCACTCAGGACACAGCCAG	<u>60</u>	30			
KNQ1-7F	TTGCAGACATAGGGTGCACACGTGC	<u>61</u>	25	1589BP	7F/7R	16
KNQ1-7R	AACAGGAGCGACGTCGCTAAGCTAG	<u>62</u>	25			

Table 7: MYO7A primers

primer name	primer sequence	SEQ ID NO.	bases	product size	primer pair	exons
MYO7A-1F	AGCACATCAGTGATTAAGTCAGG	<u>63</u>	23	822 bp	1F/1R	1
MYO7A-1R	GATTGATGGACAACATGCTCCT	<u>64</u>	23			
MYO7A-2F	TTGGGAATCTCTGAATGACAGTG	<u>65</u>	23	434 bp	2F/2R	2
MYO7A-2R	GGTTTGGAAAGCCTAGGCAGGAA	<u>66</u>	22			
MYO7A-3F	GAGAGGCCCTGGCTCTCTGA	<u>67</u>	22	628 bp	3F/3R	3
MYO7A-3R	TCTCTAACACCATGCAGAGTGG	<u>68</u>	22			
MYO7A-4F8	CTGATGTCCAGATTCTGCTAGT	<u>69</u>	23	2863bp	4F8/4R8	4
MYO7A-4R8	ACCTCCAGCATTATTATGCCATG	<u>70</u>	25			
MYO7A-5F	AGAAGGAAATCTAGGCTTAGAGACTCCACCTCCC	<u>71</u>	34	7707 bp	5F/5R	5 to 14
MYO7A-5R	GCATATGATTCACCTATATGAGGTACCTAGAAT	<u>72</u>	34			
MYO7A-6F	TGGATGTGGTGGAACTAGGTGG	<u>73</u>	22	488 bp	6F/6R	15
MYO7A-6R	AACCGATCCCTGACCGGTTCTG	<u>74</u>	22			
MYO7A-7F1a	AGAGGTGGTAACTTGGAAGTCCCTGG	<u>75</u>	26	7573bp	7F1a/7R1a	16 to 21
MYO7A-7R1a	GGTATGTGCACTCCTCAGAGCAGGCATA	<u>76</u>	28			
MYO7A-7F1d	TGGTCAGATGGATAGATGGCATCACCTC	<u>77</u>	28	4102 bp	7F1d/7R1a2	16 to 18
MYO7A-7R1a2	ATCACATCTTGCTGATGAGGAAATGCAGG	<u>78</u>	29			
MYO7A-7F1e	TCACAGTCTGGTGGCATAGTACCTAAATTG	<u>79</u>	30	4128 bp	7F1e/7R1a1	16 to 18
MYO7A-7R1a1	CTCCCAGGTTGATGATGATCTCAAACAC	<u>80</u>	28			
MYO7A-7F21a	TGCAGCTCCTGATCTAGGAT	<u>81</u>	20	591 bp	7F21a/7R21a	21
MYO7A-7R21a	AGAGCAGGCATAACTGCAG	<u>82</u>	21			
MYO7A-7F21b	ATTAGAGATCTCAGACAGGGTG	<u>83</u>	22	898bp	7F21b/7R21b	21
MYO7A-7R21b	AACTGGGATGACTTTGATAGG	<u>84</u>	22			
MYO7A-7F2a	ACCTCAGTCACTCTGGGAATCTCTG	<u>85</u>	26	3361bp	7F2a/7R2a	22 to 26
MYO7A-7R2a	TAGAAGTGTATCCCTCTCAGCTGTG	<u>86</u>	26			
MYO7A-8F	TGCAGGGTATCGAGGAGGTGGC	<u>87</u>	22	620 bp	8F/8R	27
MYO7A-8R	TGCAATATCTCCAAGGGATGCC	<u>88</u>	22			
MYO7A-9F1	GGCCCCCTTAAGTATTACACATTACAGAAATA	<u>89</u>	32	11,772bp	9F1/9R3	28 to 35
MYO7A-9R3	GTTGAAACTGATCTCCAGTGTGGCAGTGG	<u>90</u>	32			
MYO7A-10F	CGAGGTGGAAGGAGTCTGGAGGCCGCTCACAA	<u>91</u>	34	8018 bp	10F/10R	36 to 44

Appl. No. : 10/786,518
 Filed : February 4, 2004

MYO7A-10R	AGACACATAATAGAGGCTAACATGCAAGCTCC	<u>92</u>	34			
MYO7A-11F	GGCCATGCACTCCAAC TGCCAAC TGCTGAGTCT	<u>93</u>	33	4555 bp	11F/11R	45 to 49
MYO7A-11R	TCACCTCCCAGCCTGATGTCCAGCACTTCCCTCC	<u>94</u>	33			

Table 8: OTOF primers

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
OTOF-1F	TGGTAGCACATAAGCCTCTG	<u>95</u>	20	1001	1F/1R	1
OTOF-1R	ATCACAATGGCCAGTCAGTC	<u>96</u>	20			
OTOF-2F	TCCTAACATGGAAC TCA TGG	<u>97</u>	20	451	2F/2R	2
OTOF-2R	TTACCACCTCCTTCAGGAAG	<u>98</u>	20			
OTOF-3F	CCAACATCTCTGAGCACCAT	<u>99</u>	20	786	3F/3R	3
OTOF-3R	TGAGTGTCTGAGATCAGGC	<u>100</u>	19			
OTOF-4F	ACAAACAACCATCCACAGTGGG	<u>101</u>	22	3197	4F/4R	4 to 5
OTOF-4R	TCTGAGAAAGGCAGGAGATCTAG	<u>102</u>	23			
OTOF-5F	AAAGACAAGTCAGGCTTGAGCAC	<u>103</u>	24	2937	5F/5R	6 to 8
OTOF-5R	TATGAAGTCCAATACTGAACATG	<u>104</u>	23			
OTOF-6F	TGTGGTAGTGCATGCCTGTAATCC	<u>105</u>	24	6513	6F/6R	9 to 11
OTOF-6R	ATGGCTGTGTACTAACAGTCGC	<u>106</u>	24			
OTOF-7F1a	AGCTCCAGAGGACCTCAGACTCTATC	<u>107</u>	26	4152	7F1a/7R1a	12 to 25
OTOF-7R1a	TGAGGTATGACTCCTCAGGTAGACAG	<u>108</u>	26			
OTOF-7F2a	CCTGCTTCCATGGATATCCAGGCT	<u>109</u>	24	5373	7F2a/7R2a	16 to 25
OTOF-7R2a	CTCAGTCTGTAGGAGACAGGGAGGTGA	<u>110</u>	26			
OTOF-7F2e	CTGTGGAGATCGTAGACACCTCCAA	<u>111</u>	25	1791	7F2e/7R2e	16 to 18
OTOF-7R2e	ACTAGAGGTGGCTCCTGTCCTTGTC	<u>112</u>	25			
OTOF-7F2f	TAACTACACGCTGCTGGATGAGC ATC	<u>113</u>	26	1784	7F2f/7R2d	16 to 18
OTOF-7R2d	AGACCAGCTTGTGTGTTCCAGGGAAG	<u>114</u>	27			
OTOF-7F2e	CTGTGGAGATCGTAGACACCTCCAA	<u>115</u>	25	3315	7F2e/7R2i	16 to 20
OTOF-7R2i	CTCTGTAGATTCTCCTCATCTGCC	<u>116</u>	26			
OTOF-7F2f	TAACTACACGCTGCTGGATGAGC ATC	<u>117</u>	26	3404	7F2f/7R2i	16 to 20
OTOF-7F2m	TGATCAACAGGGAGGGAGGCATT	<u>118</u>	23	955	7F2m/7R2m	19 to 20
OTOF-7R2m	CTGCCCCCTCCAGCACCTTA	<u>119</u>	20			
OTOF-7F2n	CCTAGCGAGAGCTCCAG	<u>120</u>	18	542	7F2n/7R2n	19 to 20
OTOF-7R2n	GACAGCTCGGGCCATGAC	<u>121</u>	18			
OTOF-7F3f1	TGGCAGATGAGGAAGAACATACAGAGC	<u>122</u>	28	2838	7F3f1/7R3a1	21 to 25
OTOF-7R3a1	TTACCACAGCGCCATGAGTTGTTGAAG	<u>123</u>	28			
OTOF-7R3b1	ACATGAGGTCCCTACCTCTAGTCCAG	<u>124</u>	28	2697	7F3f1/7R3b1	21 to 25
OTOF-7F-A	CTGTGGAGATCGTAGACACCTCCAACCCCTGAGCT	<u>125</u>	34	16,256	7F-A/7R-A	16 to 39

Appl. No. : 10/786,518
 Filed : February 4, 2004

OTOF-7R-A	CAGATAGCCTCTACCTCACTGGGATTTGGACA	<u>126</u>	34			
OTOF-8F5	TAAGGACCAAACGAGATCACAGGTGTGGA	<u>127</u>	29	10127	8F5/8R6	26 to 39
OTOF-8R6	AGCCTCTACTCACTGGGATTTGGACA	<u>128</u>	29			
OTOF-8R7	CGAGTCACTAGAAGTAGGATCTGGTTGT	<u>129</u>	30	10181	8F5/8R7	26 to 39
OTOF-8R4	GGTTGTTACCTCACTGGGATTTGGACA	<u>130</u>	30	10128	8F5/8R4	26 to 39
OTOF-9F1	GTAGACAGGTGATGGCATAGAGGTTCT	<u>131</u>	28	7106	9F1/9R1	40 to 47
OTOF-9R1	TGGTACTGAATCTGCCAGCCTAGAGAAC	<u>132</u>	28			
OTOF-9F2	AGGCACCTCCCAGAGAACGAGAACATTG	<u>133</u>	28	7759	9F2/9R9	40 to 47
OTOF-9R9	TGTGGCTGAATCTTTAAAGAGGTCAGG	<u>134</u>	29			

Table 9: SLC26A4 sequences

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
SLC-1F	TCAGAGAATTGATCAGGGTTCTC	<u>135</u>	25	3665	1F/1R	1 to 3
SLC-1R	TAAGCAACCATCTGTACAGACC	<u>136</u>	23			
SLC-2F2	TGGAACCATTGTAAGTTGAGGACTT	<u>137</u>	25	3225	2F2/2R4	4 to 6
SLC-2R4	GAGATGAGGTCTCACGCTCTAAACT	<u>138</u>	25			
SLC-3F	ATCAACTGGGAGTTTCAGGTTATCAGCC	<u>139</u>	29	7618	3F/3R	7 to 10
SLC-3R	AAGGCAAATTGCTCTGCTAACGCTCGGTG	<u>140</u>	28			
SLC-4F	AATGAGACCATGTCTACAAGTACGAAGTG	<u>141</u>	30	11306	4F/4R	11 to 18
SLC-4R	TTTGTTCACTCTTACCTAGGTGAGAGCCTG	<u>142</u>	30			
SLC-5F4	GATCGTCCACAAGGTTGACTACGACCAGT	<u>143</u>	28	9069	5F4/5R6	19 to 21
SLC-5R6	TCATTGATTCTCACCTCACAGATCTAACG	<u>144</u>	29			

Table 10: USH2A sequences

primer name	primer sequence	SEQ ID NO:	bases	product size	primer pair	exons
USH2A-1F	TAGGATAAGGTGACTGCTACTT	<u>145</u>	23	5085	1F/1R	1 to 3
USH2A-1R	GAAGACAAATCCTGTGTTAACCA	<u>146</u>	25			
USH2A-2F	AACACATGGAGATATCACTGAGC	<u>147</u>	23	699	2F/2R	4
USH2A-2R	CCTAAATCCAATGACAAGTGTCCCT	<u>148</u>	25			
USH2A-3F1	CTTAAGTCCTACAGTGTCCATGGAGATA	<u>149</u>	28	7298	3F1/3R1	5 to 9
USH2A-3R1	CATCAGTGATGTGTTAAAGGTTATATT	<u>150</u>	28			
USH2A-4F	TCACTGATATGTGCTTACTTCTGG	<u>151</u>	25	3302	4F/4R	10 to 11
USH2A-4R	AGGATTTCCTGGCAAATGCAGTCTTC	<u>152</u>	26			
USH2A-5F	GTCTTGTACCTAATGAGCAAATTATCT	<u>153</u>	27	4954	5F/5R	12 to 13
USH2A-5R	GCATTGTATGGATATTCAACTCAAATT	<u>154</u>	27			
USH2A-6F1	GAATTAGTGCCTTGGTAGA	<u>155</u>	19	378	6F1/6R	14
USH2A-6F2	GTATTGGGAATTAGTGCCTT	<u>156</u>	20	386	6F2/6R	14
USH2A-6R	CAGAAGTTATTGCTTGCAACT	<u>157</u>	22			

Appl. No. : 10/786,518
Filed : February 4, 2004

USH2A-7F	CTCTACAATGCTATTGGTAGGTGTAACCTTA	<u>158</u>	30	10458	7F/7R	15 to 16
USH2A-7R	CACAAACAGCATTATCCTCAATGTCAAAGA	<u>159</u>	30			
USH2A-8F	AGCAGTTAGCAATGATTCTTACCAACTTGTG	<u>160</u>	32	10312	8F/8R	17 to 20
USH2A-8R	CCTGGAGTCACGCTACAACTAATTACATTCT	<u>161</u>	32			
USH2A-9F	TTCCTAGAGCCATACAGATACTTG	<u>162</u>	24	1826	9F/9R	21
USH2A-9R	GCTGAATGGAAACGGATGCTATT	<u>163</u>	23			